

SEQUENCE LISTING

<110> BRAIN AG
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<120> ISOLATION AND CLONING OF DNA FROM UNCULTIVATED ORGANISMS

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<213> Crenarchaeote

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Tyr Ile Gly Glu Lys Lys Ser Val Phe Leu Lys Phe Tyr Asn Pro Glu
 35 40 45

Asp Ser Gln Ile Tyr Phe Trp Ser Glu Ser Phe Ile Glu Asn His Ile
 50 55 60

Asn Lys His Gln Pro Tyr Cys Phe Val Lys Glu Leu Tyr Ser Asp Gln
 65 70 75 80

Val Lys Thr Ile Val Ser Lys Glu Pro His Arg Phe Arg Leu Glu Lys
 85 90 95

Ile Lys Lys Met Asp Asp Ile Glu Asp Lys Glu Ile Ser Val Phe Lys
100 105 110

Ile Ile Ala Pro Asp Pro Leu Ser Ile Gly Gly Thr Asp Ser Ser Phe
115 120 125

Arg Glu Lys Val Thr Ser Trp Glu Ala Asp Ile Lys Tyr His Glu Ser
130 135 140

Tyr Leu Phe Asp Leu Gly Leu Ile Pro Gly Ala Phe Tyr Asn Arg Ile
145 150 155 160

Gly Asn Asn Leu Val Phe His Glu Phe Pro Met Pro Glu Lys Val Asp
165 170 175

Glu Tyr Leu Asp Asn Leu Ile Lys Pro Asn Phe Lys Glu Asn Glu Pro
180 185 190

Lys Ser Ser Glu Tyr Asn Glu Phe Leu Ile Lys Trp Ser Arg Leu Leu
195 200 205

Asn Gln Pro Ile Pro Asp Ile Lys Arg Ile Ser Leu Asp Ile Glu Val
210 215 220

Asp Ser Glu Glu Gly Arg Met Pro Thr Ala Arg Asp His Asp Lys Val
225 230 235 240

Ile Thr Ala Val Gly Leu Ser Ala Ser Asp Gly Phe Arg Lys Val Phe
245 250 255

Val Leu Arg Lys Asp Pro Asn Phe Asp Pro Ser Lys Leu Asp Ser Thr
260 265 270

Thr Val Glu Leu Cys Asp Ser Glu Lys Asp Met Ile Leu Lys Val Phe
275 280 285

Ala Ile Ile Gln Asn Tyr Pro Ile Val Leu Thr Phe Asn Gly Asp Asp
290 295 300

Phe Asp Leu Pro Tyr Leu Tyr Ala Arg Ser Gln Asp Pro Ser Ile Asp
305 310 315 320

Pro Val His Lys Lys Pro Ile Ser Lys Glu Leu Val Pro Ile Leu Val
 325 330 335

Lys Lys Asp Ser Phe Ile Lys Arg Gly Ile Gln Ala Asp Pro Val Ser
 340 345 350

Leu Lys His Gly Ile His Ile Asp Leu Phe Arg Thr Phe Gln Asn Lys
 355 360 365

Ser Val Gln Asn Tyr Ala Phe Ser His Lys Tyr Ser Glu Phe Thr Leu
 370 375 380

Asn Ala Ile Cys Glu Ala Leu Leu Asn Glu Ser Lys Ile Asp Phe Asp
 385 390 395 400

Glu Ser Ile Gly Asp Leu Pro Leu Glu Lys Leu Ala Glu Tyr Cys Leu
 405 410 415

Lys Asp Ala Asp Leu Thr Phe Arg Leu Thr Ser Phe Asn Asp Asn Leu
 420 425 430

Leu Ile Lys Leu Leu Ile Ile Ile Ser Arg Ile Ser Arg Met Ser Ile
 435 440 445

Glu Asp Ile Thr Arg Phe Gly Val Asn Gln Trp Ile Arg Ser Met Met
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Phe Phe Glu His Arg Gln Gln Asn Ile Ile Ile Pro Arg Lys Asp Glu
 465 470 475 480

Leu Gln Lys Lys Gly Thr Ser Ser Thr Val Ala Ile Ile Lys Glu Lys
 485 490 495

Lys Tyr Arg Gly Gly Leu Val Val Glu Pro Val Leu Gly Ile His Phe
 500 505 510

Asn Val Ile Val Val Asp Phe Ala Ser Leu Tyr Pro Ser Ile Ile Lys
 515 520 525

Val His Asn Leu Ser Tyr Glu Thr Val Asn Cys Pro His Glu Asn Cys
 530 535 540

Arg Arg Asp Pro Ser Thr His Ile Glu Gln Thr Asn His Trp Val Cys
 545 550 555 560

Lys Glu Lys Gln Gly Met Thr Ser Ile Leu Ile Gly Thr Leu Arg Asp
 565 570 575

Leu Arg Val Asn Tyr Tyr Lys Tyr Leu Ser Lys Asp Asn Ser Leu Asp
 580 585 590

Lys Glu Asp Lys Gln Leu Tyr Ser Val Ile Ser Gln Ala Ile Lys Val
 595 600 605

Ile Leu Asn Ala Thr Tyr Gly Val Met Gly Ala Glu Ile Phe Pro Leu
 610 615 620

Tyr Cys Leu Pro Val Ala Glu Ala Thr Ala Ala Val Gly Arg Met Thr
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Thr Thr Lys Thr Ile Glu Lys Cys Asn Glu Glu Lys Ile Glu Val Ile
 645 650 655

Tyr Gly Asp Thr Asp Ser Leu Phe Leu Lys Asn Pro Ser Lys Glu Gly
 660 665 670

Leu Ser Gly Ile Ser Ser Trp Ser Lys Lys Glu Leu Gly Ile Asp Leu
 675 680 685

Glu Ile Asp Lys Arg Tyr Arg Tyr Val Val Phe Ser Glu Leu Lys Lys
 690 695 700

Asn Tyr Leu Gly Val Leu Glu Asp Gly Thr Val Asp Val Lys Gly Leu
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Thr Gly Lys Lys Ser His Thr Pro Pro Ile Ile Arg Gln Ala Phe Tyr
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 740 745 750

Arg Ala Lys Glu Lys Ile Lys Lys Ile Val Gln Ser Ile Ala Glu Asn
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Thr Met Asp Phe Ile Lys Ile Leu Gly Tyr Asn Ile Arg Tyr Ile Lys
35 40 45

Ile Asp Gln Val Lys Ser Asn Glu Thr Ile Ile Leu Leu His Gly Ile
50 55 60

Gly Ala Ser Ala Glu Arg Trp Ser Glu Leu Val Pro Phe Leu Tyr Asn
65 70 75 80

Cys Asn Ile Ile Ile Pro Asp Ile Ile Gly Phe Gly Tyr Ser Glu Lys
85 90 95

Pro Arg Ile Glu Tyr Asn Ile Asp Leu Phe Val Lys Phe Leu Asp Glu
100 105 110

Leu Phe Leu Lys Leu Glu Ile Lys Asn Pro Ile Ile Met Gly Ser Ser
115 120 125

Phe Gly Gly Gln Leu Ile Leu Glu Tyr Tyr Phe Arg His Lys Asp Phe
130 135 140

Phe Lys Lys Met Ile Leu Val Ser Pro Ala Gly Thr Gln Glu Arg Pro
145 150 155 160

Thr Leu Ala Leu Arg Gln Tyr Thr Tyr Ser Cys Leu Tyr Pro Thr Arg
165 170 175

Glu Asn Thr Glu Arg Ala Phe Lys Met Met Ser His Phe Asn His Thr
180 185 190

Val Lys Asp Ser Met Ile Lys Asp Phe Ile Asn Arg Met Lys Gln Pro
195 200 205

Asn Ala Lys His Ser Phe Val Ser Thr Leu Leu Ala Leu Arg Lys Asn
210 215 220

Ser Asp Leu Gln Asp Asn Leu Arg Glu Ile Lys Ile Pro Thr Leu Val
225 230 235 240

Ile Trp Gly Lys Glu Asp Asn Thr Ile Pro Val Glu Asn Ile Glu Tyr
245 250 255

Phe Arg Gly Ile Pro Phe Val Lys Thr Cys Ile Met Ser Asp Cys Gly
260 265 270

His Val Pro Phe Val Glu Lys Pro Leu Glu Phe Tyr Lys Ile Val Lys
275 280 285

Glu Phe Ile Asp Ser
290

<210> 6

<211> 318

<212> DNA

<213> Crenarchaeote

<400> 6

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caatacaatg cgtttcagaa tatgttttca tctttgcaag gatttacaaa ttataatgcc	180
atgttttaaaa ccaccgtaca gacgggtggc aggatctcaa ttcccgaagc agaaagaaat	240
gctttggggg ttgaagaggg tgatctagtc caggttataa ttataccggt gacaaggaaa	300
aagaaaaaca caagttaa	318

<210> 7

<211> 105

<212> PRT

<213> Crenarchaeote

<400> 7

Met Asn Gln Ser Thr Ser Met Ser Asn Glu Asn Glu Glu Asn Lys Asp
1 5 10 15

Ile Asp Phe Lys Lys Ser Ile Glu Lys Ala Ala Glu Phe Gln Gln Asp
20 25 30

Leu Leu Arg Gln Phe Ser Thr Ile Gln Tyr Asn Ala Phe Gln Asn Met
35 40 45

Phe Ser Ser Leu Gln Gly Phe Thr Asn Tyr Asn Ala Met Phe Lys Thr
50 55 60

Thr Val Gln Thr Gly Gly Arg Ile Ser Ile Pro Glu Ala Glu Arg Asn
65 70 75 80

Ala Leu Gly Ile Glu Glu Gly Asp Leu Val Gln Val Ile Ile Ile Pro
85 90 95

Leu Thr Arg Lys Lys Lys Asn Thr Ser
100 105

<210> 8

<211> 1086

<212> DNA

<213> Crenarchaeote

<400> 8

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gaaaacataa agactggatc aagcgaatat gaggtaataa gggaacaac cctcttcggt 180

ttactacatt ataaacccct aaagcaacaa actttcaagt accctttgtt gattgtttat	240
gcattaataa acaaatcata ttttttggat ctgcagaacg acaaaagttg gataaggaac	300
ctgctagagc agggcataaa tgtctatctg attgactgga aacccccgtc aaaactggat	360
aaatacatca ctgttgatga ttatgtcaat ttgtttattt atgagtgtgt agaatacata	420
aaaaacatag aaaacattga tcagatttca ttacaaggat attgcatggg ggggtacaatg	480
tccttgatgt acacttcgct atatcaaaaa aacattaaaa atctagtcac cattgctcca	540
attgttgatg ccgagaaaaga caaatccgta ataaaaaaca tggctgagca catggatatt	600
gacaaagtac tgtcctatca cgaaaacttt ccatatgaat tactgtatct ggtttatgca	660
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aaaaacaaga tgattgtggg tgaaaacaag ataaatttgt caaacattaa ggttcccgtt	900
cttaatgttg tagctgaatt tgaccacctt gtaacgtctg acagcagtag ctccctaaac	960
aacctaatTT caagtcagga taaaagcctg atgaaatttc caacagggca tgtagggcta	1020
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<210> 9

<211> 361

<212> PRT

<213> Crenarchaeote

<400> 9

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1				5					10					15	

Tyr	Phe	Thr	Phe	Val	Arg	Asp	Pro	Asp	Asn	Ile	Ser	Lys	Leu	Lys	Glu
			20					25					30		

Ile	Arg	Lys	Lys	Leu	Ser	Asn	Ile	Glu	Asn	Ile	Lys	Thr	Gly	Ser	Ser
		35					40					45			

Glu Tyr Glu Val Ile Arg Glu Thr Thr Leu Phe Arg Leu Leu His Tyr
 50 55 60

Lys Pro Leu Lys Gln Gln Thr Phe Lys Tyr Pro Leu Leu Ile Val Tyr
 65 70 75 80

Ala Leu Ile Asn Lys Ser Tyr Ile Leu Asp Leu Gln Asn Asp Lys Ser
 85 90 95

Trp Ile Arg Asn Leu Leu Glu Gln Gly Ile Asn Val Tyr Leu Ile Asp
 100 105 110

Trp Lys Pro Pro Ser Lys Leu Asp Lys Tyr Ile Thr Val Asp Asp Tyr
 115 120 125

Val Asn Leu Phe Ile Tyr Glu Cys Val Glu Tyr Ile Lys Asn Ile Glu
 130 135 140

Asn Ile Asp Gln Ile Ser Leu Gln Gly Tyr Cys Met Gly Gly Thr Met
 145 150 155 160

Ser Leu Met Tyr Thr Ser Leu Tyr Gln Lys Asn Ile Lys Asn Leu Val
 165 170 175

Thr Ile Ala Pro Ile Val Asp Ala Glu Lys Asp Lys Ser Val Ile Lys
 180 185 190

Asn Met Ala Glu His Met Asp Ile Asp Lys Val Leu Ser Tyr His Glu
 195 200 205

Asn Phe Pro Tyr Glu Leu Leu Tyr Leu Val Tyr Ala Ser Leu Lys Pro
 210 215 220

Phe Lys Gln Gly Val Asn Lys Tyr Tyr Asn Leu Phe Lys Asn Phe Glu
 225 230 235 240

Asp Glu Ser Phe Val Gln Asn Phe Leu Arg Ile Glu Lys Trp Leu Tyr
 245 250 255

Asp Thr Pro Pro Ile Ala Gly Glu Thr Phe Arg Gln Trp Val Lys Asp
 260 265 270

Ile Tyr Gln Gln Asn Leu Phe Ala Lys Asn Lys Met Ile Val Gly Glu
 275 280 285

Asn Lys Ile Asn Leu Ser Asn Ile Lys Val Pro Val Leu Asn Val Val
 290 295 300

Ala Glu Phe Asp His Leu Val Thr Ser Asp Ser Ser Ser Ser Leu Asn
 305 310 315 320

Asn Leu Ile Ser Ser Gln Asp Lys Ser Leu Met Lys Phe Pro Thr Gly
 325 330 335

His Val Gly Leu Ile Ala Ser Asn Phe Ser Gln Lys Asn Val Leu Pro
 340 345 350

Lys Ile Gly Lys Trp Ile Gln Thr His
 355 360

<210> 10

<211> 582

<212> DNA

<213> Crenarchaeote

<400> 10

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cttacaacct tggggccatt cacctcttta atgaatgatc caaaccttaa tttaaataca	180
ttaaaggaac acggtaatTT gttactgaga tatcagtcac ttctcaacct atacttttcc	240
cgtatgataa atgcttattt gttggccgta aacaaggatc cgtctgctat agatgaaaaa	300
aaccccgacg atattaggaa aataatcata aatacttttg aggatgtggt ctcgtcaatg	360
ttgcagtcaa cagacttttc aatcaattat aacaatttat tgaattccag cattgatgtc	420
atcaaaaagt atcaaaaaat ttacgattca aatgccggtt tgtttagggtc acaacaacaa	480
ctgtcaaaaag aagaaaaaga cctgttattt tataatctct atgaaatcaa aaaaatatca	540
ttggaaatca aaaaaaaatt aaatgagaaa aaaaatgaat aa	582

<210> 11

<211> 193

<212> PRT

<213> Crenarchaeote

<400> 11

Met Gln Leu Glu Asn Asn Asn Ile Gly Glu Glu Lys Asn Ser Lys Asn
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Thr Leu Ser Glu Glu Ala Gly Leu Gln Ser Val Phe Glu Asn Phe Ile
20 25 30

Lys Gln Leu Thr Glu Leu Asn Ser Leu Thr Thr Leu Gly Pro Phe Thr
35 40 45

Ser Leu Met Asn Asp Pro Asn Leu Asn Leu Asn Thr Leu Lys Glu His
50 55 60

Gly Asn Leu Leu Leu Arg Tyr Gln Ser Phe Leu Asn Leu Tyr Phe Ser
65 70 75 80

Arg Met Ile Asn Ala Tyr Leu Leu Ala Val Asn Lys Val Ser Ser Ala
85 90 95

Ile Asp Glu Lys Asn Pro Asp Asp Ile Arg Lys Ile Ile Ile Asn Thr
100 105 110

Phe Glu Asp Val Phe Ser Ser Met Leu Gln Ser Thr Asp Phe Ser Ile
115 120 125

Asn Tyr Asn Asn Leu Leu Asn Ser Ser Ile Asp Val Ile Lys Ser Tyr
130 135 140

Gln Lys Ile Tyr Asp Ser Asn Ala Val Leu Phe Arg Ser Gln Gln Gln
145 150 155 160

Leu Ser Lys Glu Glu Lys Asp Leu Leu Phe Tyr Asn Leu Tyr Glu Ile
165 170 175

Lys Lys Ile Ser Leu Glu Ile Lys Lys Lys Leu Asn Glu Lys Lys Asn
180 185 190

Glu

<210> 12

<211> 438

<212> DNA

<213> Crenarchaeote

<400> 12

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aagagggtta tgtgggggac aatagaaaga atccatccac aaacgaatgg ttttggcaaa	180
tgtctgcctt ggctcattgt ttctatatat ggtttcgcca tagataatat ttgggtaatt	240
actcatgtgc ctgtatacaa aaatagacaa ccatctctac ctatatataa attttttgac	300
aagggtcagg ttctttttct cttttttttg caaattatgc cgggccaccc agaacaac	360
cccgctttg ggcccgaat gattactggg caacccaaat ctagcgcccc acgcccagga	420
gtgcaagtga caatgtga	438

<210> 13

<211> 145

<212> PRT

<213> Crenarchaeote

<400> 13

Met Pro Thr Ser Ser Asp Val Leu Tyr Met Ser Lys Pro Ala Val Val	
1 5 10 15	

Cys Ile His Ser Phe Asp Met Val Gly Gly Tyr Ala His Thr Gln Lys	
20 25 30	

Leu Arg Cys Cys Ile Ser Leu Gly Lys Arg Val Met Trp Gly Thr Ile
 35 40 45

Glu Arg Ile His Pro Gln Thr Asn Gly Phe Gly Lys Cys Leu Pro Trp
 50 55 60

Leu Ile Val Ser Ile Tyr Gly Phe Ala Ile Asp Asn Ile Trp Val Ile
 65 70 75 80

Thr His Val Pro Val Tyr Lys Asn Arg Gln Pro Ser Leu Pro Ile Tyr
 85 90 95

Lys Phe Phe Asp Lys Gly Gln Val Leu Phe Leu Leu Phe Leu Gln Ile
 100 105 110

Met Pro Gly His Pro Glu Thr Asn Pro Ala Phe Gly Pro Ala Met Ile
 115 120 125

Thr Gly Gln Pro Lys Ser Ser Ala Pro Arg Pro Gly Val Gln Val Thr
 130 135 140

Met
 145

<210> 14

<211> 915

<212> DNA

<213> Crenarchaeote

<400> 14
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 gcaaaaaatc tgaccaacag aattgccatt gtaggaactg gaatagataa aaatatcttt 360
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ggaaggatag aaaaatttca tggactggag gaaatttggga ctgcaataaa aacactcaga 480
 ccagaatcta attttgtaat ggttggggcgc atacccttg ataaggctgc aaaactacgt 540
 aatgcgggta tagatcacag aggctttgtc tccgaggaag aaaagattag cctttattct 600
 aaatctaaag tctttatttt tccatcatcc agagaggggtt ttggcattgc tgtggctgag 660
 gccttagttt cgtgtgttcc cactgttgcc tggaaactcc ccgtttttga agaactatac 720
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 gaagagtgcg taaaaatgct aaataaatat ggcataatca aaaaggcgac tgaaggaaaa 840
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<210> 15

<211> 304

<212> PRT

<213> Crenarchaeote

<400> 15

Met Asp Ser Trp Gly Glu Ser Asn Ile Val Leu Trp Leu Leu Leu Arg
 1 5 10 15

Leu Phe Lys Pro Lys Thr Lys Ile Phe Val Val Phe His His His Glu
 20 25 30

Pro Arg Ile Ser Ile Cys Lys Asn Tyr Phe Glu Phe Leu Tyr Asn Tyr
 35 40 45

Leu Ile Gln Lys Ala Thr Ala Val Met Leu Lys Asp Ser Asp Met Ile
 50 55 60

Leu Thr Val Ser Gln Ala Ser Lys His Glu Leu Asn Thr Val Tyr Gly
 65 70 75 80

Ile Gly Val Ser Lys Ile Asn Asn Leu Lys Glu Thr Ala Asn Lys Lys
 85 90 95

Thr Arg Glu Leu Ala Lys Asn Leu Thr Asn Arg Ile Ala Ile Val Gly
100 105 110

Thr Gly Ile Asp Lys Asn Ile Phe Leu Lys Asp Ser Asn Arg Gly Val
115 120 125

Ile Asn Asn Lys Lys Asp Ile Asp Phe Leu Cys Ile Gly Arg Ile Glu
130 135 140

Lys Phe His Gly Leu Glu Glu Ile Trp Thr Ala Ile Lys Thr Leu Arg
145 150 155 160

Pro Glu Ser Asn Phe Val Met Val Gly Arg Ile Pro Pro Asp Lys Ala
165 170 175

Ala Lys Leu Arg Asn Ala Gly Ile Asp His Arg Gly Phe Val Ser Glu
180 185 190

Glu Glu Lys Ile Ser Leu Tyr Ser Lys Ser Lys Val Phe Ile Phe Pro
195 200 205

Ser Ser Arg Glu Gly Phe Gly Ile Ala Val Ala Glu Ala Leu Val Ser
210 215 220

Cys Val Pro Thr Val Ala Trp Lys Leu Pro Val Phe Glu Glu Leu Tyr
225 230 235 240

Leu Lys Asn Gly Asn Thr Asn Ile Lys Leu Ile Glu Tyr Gly Glu Thr
245 250 255

Thr Leu Phe Ala Glu Glu Cys Val Lys Met Leu Asn Lys Tyr Gly Ile
260 265 270

Ile Lys Lys Ala Thr Glu Gly Lys Lys Val Ser Phe Gln Leu Pro Asn
275 280 285

Trp Gln Thr Val Ala Lys Asn Val Met Thr Thr Ile Glu Ser Val Thr
290 295 300

<210> 16

<211> 1692

<212> DNA

<213> Crenarchaeote

<400> 16

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aaaatgctat cctgtatgaa aaaccgggggt ccggatggca tgggtttgtc tacagagaat	120
caaatagttt attctgatac ctttgataat ccattgtttt cacaggtaga ggggcatgac	180
gttttaggtc acagtcgttt ggcaatagtt ggtggctcct gtggtcagca gccgtttgtg	240
agttgtgata aaaaaactcat tctggagcat aatgggtgaaa tatataacta taaagaaatc	300
agaaagaacc tttctgcaca tcacactttt actacctcga ctgatagtga agttattgtt	360
caccttcttg aagaccatta tcaaaacact aaaggcgatc taatcgaagc tataaggaga	420
accgttacct agcttgatgg aatttatgtt ttggcgatta gagagcagtc cacaggagat	480
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ttcattgcat ttgcatcaga aagaaaagcc ttatggaaaa ttgctatgtc cgaccaaadc	600
aaaagacttt tgccaggcta tgctcttgct atttcgcgga aggaagggtc ctccaatttc	660
aagactacat tgtttccgat ttctgtaaat acaaaaaaat caatatgtga gaaatattca	720
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gttaaattgg ctacacgaaca gggaatacgg gtaatgctta caggtcaggg ggacagacgaa	1140
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cagggatacc tgatagagga cattaagtta ctttacaaag aaacactgga aagagaggac	1260
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gacacggtac tgagaataga tccgcgactc aatattcaaa acaatggcaa taactatgac	1380

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<210> 17

<211> 563

<212> PRT

<213> Crenarchaeote

<400> 17

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Pro Leu Ile Gly Lys Met Leu Ser Cys Met Lys Asn Arg Gly Pro Asp
 20 25 30

Gly Met Gly Leu Ser Thr Glu Asn Gln Ile Val Tyr Ser Asp Thr Phe
 35 40 45

Asp Asn Pro Leu Phe Ser Gln Val Glu Gly His Asp Val Leu Gly His
 50 55 60

Ser Arg Leu Ala Ile Val Gly Gly Ser Cys Gly Gln Gln Pro Phe Val
 65 70 75 80

Ser Cys Asp Lys Lys Leu Ile Leu Glu His Asn Gly Glu Ile Tyr Asn
 85 90 95

Tyr Lys Glu Ile Arg Lys Asn Leu Ser Ala His His Thr Phe Thr Thr
 100 105 110

Ser Thr Asp Ser Glu Val Ile Val His Leu Leu Glu Asp His Tyr Gln
 115 120 125

Asn Thr Lys Gly Asp Leu Ile Glu Ala Ile Arg Arg Thr Val Thr Gln
 130 135 140

Leu Asp Gly Ile Tyr Val Leu Ala Ile Arg Glu Gln Ser Thr Gly Asp
 145 150 155 160

Ile Val Leu Val Arg Asp Gly Ile Gly Val Arg Gln Ile Tyr Tyr Gly
 165 170 175

Glu Ser Ser Asp Phe Ile Ala Phe Ala Ser Glu Arg Lys Ala Leu Trp
 180 185 190

Lys Ile Ala Met Ser Asp Gln Ile Lys Arg Leu Leu Pro Gly Tyr Ala
 195 200 205

Leu Val Ile Ser Arg Lys Glu Gly Ser Ser Asn Phe Lys Thr Thr Leu
 210 215 220

Phe Pro Ile Ser Val Asn Thr Lys Lys Ser Ile Cys Glu Lys Tyr Ser
 225 230 235 240

Ile Leu Tyr Thr Asp Ile Asp Ser Ala Val Asn Ala Tyr Gly Asp Thr
 245 250 255

Leu Val Glu Ser Met Arg Lys Arg Val Ser Asp Phe Lys Lys Ile Gly
 260 265 270

Ile Val Phe Ser Gly Gly Ile Asp Ser Val Ile Val Ala Tyr Leu Ala
 275 280 285

Lys Gln Met Ala Pro Glu Val Ile Cys Tyr Thr Ser Gly Ile Lys Gly
 290 295 300

Ser Ser Asp Ile Leu Asn Ser Leu Glu Ile Ala Glu Lys Leu Asp Leu
 305 310 315 320

Lys Leu Glu Ile Glu Gln Met Thr Glu Ser Asp Val Glu Ser Thr Ile
 325 330 335

Pro Lys Ile Ile Ser Ile Ile Glu Asp Asp Asn Met Gly Gln Val Glu
 340 345 350

Val Ala Ile Pro Ile Tyr Gly Ala Val Lys Leu Ala His Glu Gln Gly
 355 360 365

Ile Arg Val Met Leu Thr Gly Gln Gly Ala Asp Glu Leu Phe Gly Gly
 370 375 380

Tyr Ser Trp Tyr Ser Lys Ile Val Lys Lys His Gly Tyr Glu Lys Ile
 385 390 395 400

Gln Gly Tyr Leu Ile Glu Asp Ile Lys Leu Leu Tyr Lys Glu Thr Leu
 405 410 415

Glu Arg Glu Asp Lys Ile Thr Met Ser Gln Ser Ile Glu Leu Arg Glu
 420 425 430

Pro Phe Leu Asp Thr Asn Leu Ile Asp Thr Val Leu Arg Ile Asp Pro
 435 440 445

Arg Leu Asn Ile Gln Asn Asn Gly Asn Asn Tyr Asp Asn Leu Gly Lys
 450 455 460

Arg Val His Arg Lys Leu Ala Glu Lys Leu Gly Ile Pro Lys Glu Ile
 465 470 475 480

Ala Tyr Arg Ile Lys Glu Ala Ala Gln His Gly Ser Gly Ile His Asn
 485 490 495

Thr Leu Asn Thr Leu Ala Met Lys Asn Gly Phe Thr Glu Ser Lys Val
 500 505 510

Asn Ser Ser Tyr Leu Asp Lys Leu Lys Lys Arg Glu Leu Ile Gly Ser
 515 520 525

Ser Gln Arg Tyr Gly His Leu Phe Glu Lys Glu Gln Ile Trp Ser Leu
 530 535 540

Glu Pro His Ile Gln Met Tyr Leu Glu Asn Ile Ser Lys Asn Ile Leu
 545 550 555 560

Pro Arg Asn

<210> 18

<211> 666

<212> DNA

<213> Crenarchaeote

<400> 18

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<210> 19

<211> 221

<212> PRT

<213> Crenarchaeote

<400> 19

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Met Leu Tyr Pro Met Glu Phe Lys Ser Thr Leu Ala Val Phe Asp Met
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Asp Gly Thr Leu Ile Asp Gly Arg Leu Ile Glu Val Leu Ser Lys Lys
          20           25           30
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Phe Gly Leu Tyr Ala Gln Val Arg His Ile Gln Ser Asp Lys Ser Ile
35 40 45

Pro Gly Tyr Val Lys Thr Gln Lys Ile Ala Ala Val Ile Arg Gly Ile
50 55 60

Glu Glu Arg Glu Ile Glu Ile Ala Leu Asp Ser Ile Pro Pro Ala Lys
65 70 75 80

Asn Ser Gln Glu Val Ile Ser Leu Leu Lys Lys Lys Gly Phe Arg Ile
85 90 95

Gly Ile Ile Thr Asp Ser Tyr Ser Val Ala Ala Gln Ala Leu Val Asn
100 105 110

Lys Leu Asp Leu Asp Phe Phe Tyr Ala Asn Glu Leu Lys Val Asp Asn
115 120 125

Gly Ile Val Thr Gly Glu Ile Asn Met Pro Leu Gly Trp Glu Lys Ile
130 135 140

Asp Cys Phe Cys Lys Asn Ser Val Cys Lys Arg Tyr His Met Glu Ile
145 150 155 160

His Ala Lys Lys Ile Cys Ala Asp Ile Lys Asn Thr Ile Ala Ile Gly
165 170 175

Asp Thr Lys Gly Asp Leu Cys Met Ile Lys Gln Ala Gly Ile Gly Ile
180 185 190

Ala Tyr Met Pro Lys Asp Lys Tyr Ile Asn Glu Thr Ile Asn Lys Val
195 200 205

Asn Thr Pro Asp Met Ile Gly Val Leu Asp Phe Ile Glu
210 215 220

<210> 20

<211> 1212

<212> DNA

<213> Crenarchaeote

<400> 20

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catcagtcta taaaggcctt tggttatgac gaactgtcaa gcatattcca agaacttgcc	180
atagtcattc cagtaaagaa cgaaaaaatc agccttcttg aaggagtatt gagcggattt	240
ccaaatgaat gtctcatcat catagtttcc aatagccaaa ggactcctgt cgacagattt	300
gccatggagg ttgaaatggg aaggcagtac tctagttttg cagacaagaa aataatgatt	360
attcaccaaa atgatcctga gctggctaata acttttaaga aaataaagta tagatccatc	420
ctcaacacca aaagtcagggt tcgtagtgga aaggctgaag gaatgataat tggaatattg	480
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tacagcaata tcagaatatc gtggcggttc aaacccaaaa tcgtaaacia ctactacaa	660
ttcccaagat ggggtagaat ttcagaatcc agtaacaaat acctgaacgc tctaatatcc	720
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ggaaatgatc atttggcagg catgatgcaa gaatctcttc tcgcaataaa caacagcaaa	1020
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cacaataatg atatgaccaa actcaacttt aagaaaaaac accttataat ggatcccata	1140
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<210> 21

<211> 403

<212> PRT

<213> Crenarchaeote

<400> 21

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Ser Ile His Ala Leu Gln Lys Ile Tyr Glu Ile Asp Ser Gly Lys Met
20 25 30

Pro Lys Phe Asn Gly Leu His Gln His Gln Ser Ile Lys Ala Phe Gly
35 40 45

Tyr Asp Glu Leu Ser Ser Ile Phe Gln Glu Leu Ala Ile Val Ile Pro
50 55 60

Val Lys Asn Glu Lys Ile Ser Leu Leu Glu Gly Val Leu Ser Gly Ile
65 70 75 80

Pro Asn Glu Cys Leu Ile Ile Ile Val Ser Asn Ser Gln Arg Thr Pro
85 90 95

Val Asp Arg Phe Ala Met Glu Val Glu Met Val Arg Gln Tyr Ser Ser
100 105 110

Phe Ala Asp Lys Lys Ile Met Ile Ile His Gln Asn Asp Pro Glu Leu
115 120 125

Ala Asn Thr Phe Lys Lys Ile Lys Tyr Arg Ser Ile Leu Asn Thr Lys
130 135 140

Ser Gln Val Arg Ser Gly Lys Ala Glu Gly Met Ile Ile Gly Ile Leu
145 150 155 160

Leu Ala Lys Met His Leu Lys Glu Tyr Ile Gly Phe Ile Asp Ser Asp
165 170 175

Asn Tyr Phe Pro Gly Ala Val Asn Glu Tyr Val Lys Ile Phe Ala Ala
180 185 190

Gly Phe Gly Met Ala Thr Thr Pro Tyr Ser Asn Ile Arg Ile Ser Trp
195 200 205

Arg Ser Lys Pro Lys Ile Val Asn Asn Ser Leu Gln Phe Pro Arg Trp
210 215 220

Gly Arg Ile Ser Glu Ser Ser Asn Lys Tyr Leu Asn Ala Leu Ile Ser
225 230 235 240

His Ile Thr Gly Phe Glu Arg Glu Ile Ile Thr Thr Gly Asn Ala Gly
245 250 255

Glu His Ala Leu Ser Met Ser Leu Ala Glu Asn Leu Asn Tyr Ser Ser
260 265 270

Gly Tyr Ser Val Glu Pro Tyr Glu Phe Ile Asn Ile Leu Glu Lys Phe
275 280 285

Gly Gly Leu Leu Pro Ser Asn Asn Pro Asp Ile Ile Glu Lys Gly Ile
290 295 300

Glu Ile Phe Gln Ile Glu Thr Arg Asn Pro His Phe His Glu Glu Lys
305 310 315 320

Gly Asn Asp His Leu Ala Gly Met Met Gln Glu Ser Leu Leu Ala Ile
325 330 335

Asn Asn Ser Lys Ile Cys Asn Thr Glu Leu Thr Arg Glu Ile Asn Asp
340 345 350

His Leu Leu Met Leu Gln Val Lys His Asn Asn Asp Met Thr Lys Leu
355 360 365

Asn Phe Lys Lys Lys His Leu Ile Met Asp Pro Ile Lys Ile Ile Pro
370 375 380

Ile Asp Lys Phe Ala Glu Phe Val Val Lys Asn Ser Lys Thr Phe Ile
385 390 395 400

Arg Ile Gly

<210> 22

<211> 1164

<212> DNA

<213> Crenarchaeote

<400> 22

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acggcgggag cccttatttt tgacagtgcc aagcaacaaa taaccccaaa acacatactt      600
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<211> 387

<212> PRT

<213> Crenarchaeote

<400> 23

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20 25 30

Ile Lys Leu Asp Ile Leu Ser Gly Thr Ser Ile Gly Gly Leu Asn Ala
35 40 45

Thr Val Ile Ala Gly Ser Lys Glu Asp Arg Pro Glu Lys Ser Leu Glu
50 55 60

Asn Phe Trp Met Glu Ile Ala Asp Thr Asn Asn Gly Asn Ile Asn Thr
65 70 75 80

Tyr Leu Asn Phe Pro Phe Phe Glu Ser Pro Phe Pro Gly Gln Ile Pro
85 90 95

Phe Pro Leu Ala Ser Glu Ser Thr Leu Ser Phe Tyr Ser Ser Ala Ile
100 105 110

Tyr Gly Asn Arg Lys Ile Phe Leu Pro Arg Trp Gly Pro Glu Asn Ile
115 120 125

Phe Lys Asp Pro Gln Tyr Phe Thr Pro Ser Lys Trp Thr Tyr Leu Tyr
130 135 140

Asp His Ser Pro Leu Val Lys Thr Leu Glu Lys Tyr Ile Asp Tyr Ser
145 150 155 160

Lys Leu Gln Pro Asn Gly Lys Pro Asn Ala Arg Leu Ile Ile Thr Ala
165 170 175

Val Asn Val Met Thr Ala Glu Pro Leu Ile Phe Asp Ser Ala Lys Gln
180 185 190

Gln Ile Thr Pro Lys His Ile Leu Ala Thr Thr Ala Tyr Pro Thr Tyr
195 200 205

Phe Phe Gln Trp Val Glu Leu Glu Lys Gly Leu Phe Ala Trp Asp Gly
210 215 220

Ser Leu Leu Ser Asn Thr Pro Leu Arg Glu Val Ile Asp Ala Ser Pro
225 230 235 240

Ala Lys Asp Lys Arg Ile Phe Leu Val Glu Asn Tyr Pro Lys Asn Ile
245 250 255

Glu Lys Leu Pro Ser Asn Leu Gln Glu Val Lys His Arg Ala Arg Asp
260 265 270

Ile Met Phe Ser Asp Lys Thr Val His Ser Ile His Met Ser Lys Ala
275 280 285

Ile Thr Leu Gln Leu Lys Leu Ile Asp Asp Leu Tyr Lys Met Leu Glu
290 295 300

Tyr Tyr Phe Asn Ser Glu Lys Ile Glu Glu Lys Glu Lys Phe Glu Lys
305 310 315 320

Ile Arg Ala Arg Tyr Lys Lys Val Ser Glu Glu His Gly Ala Glu Ile
325 330 335

Lys Gly Val Tyr Tyr Ile Thr Arg Asp Glu Pro Ser Pro Ser Leu Tyr
340 345 350

Glu Asn Ala Asp Phe Ser Lys Asn Ala Ile Lys Ala Ser Ile Asn Asp
355 360 365

Gly Glu Gln Lys Ala Asp Arg Ile Ile Lys Glu Ile Gln Thr Lys Gly
370 375 380

Lys Arg Lys
385

<210> 24

<211> 882

<212> DNA

<213> Crenarchaeote

<400> 24

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gtgggcacaa tgggtcctcc aatagcagaa tcggccttac agcaatctca actgatttgc      240
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<210> 25

<211> 293

<212> PRT

<213> Crenarchaeote

<400> 25

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Glu Gly Asn Val Ser Tyr Asn Ser Asp Gly Thr Leu Asn Arg Ala Glu
 20 25 30

Thr Lys Asn Ile Leu Gly Pro His Ser Ala Ala Ala Ser Leu Ala Ala
 35 40 45

Leu Tyr Ser Lys Val Lys His Gly Thr His Val Ser Val Gly Thr Met
 50 55 60

Gly Pro Pro Ile Ala Glu Ser Ala Leu Gln Gln Ser Gln Leu Ile Cys
 65 70 75 80

Asp Ala Asp Glu Leu His Leu Tyr Ser Asp Arg Ile Phe Ala Gly Ala
 85 90 95

Asp Thr Leu Ala Thr Ala Glu Val Leu Ile Ala Gly Ile Lys Lys Met
 100 105 110

Ala Asn Gly Gln Asp Val Asp Ile Val Phe Ser Gly His Arg Ala Ser
 115 120 125

Asp Gly Glu Thr Gly Gln Thr Gly Pro Gln Thr Ala Trp Lys Leu Gly
 130 135 140

Tyr Pro Phe Leu Gly Asn Val Ile Asp Tyr Asp Ile Asp Val Val Lys
 145 150 155 160

Arg Ile Val Arg Val Gln Arg Leu Ile Lys Ile Tyr Gly His Pro Asp
 165 170 175

Ile Ile Glu Glu Met Glu Ala Pro Leu Pro Val Phe Ile Thr Leu Asp
 180 185 190

Pro Ser Tyr Asn Pro Ser Phe Asn Thr Val Ser Gln Arg Leu Arg Leu
 195 200 205

Ala Arg Asn Leu Gln Glu Ala His Asp Arg Ser Gln Arg Tyr Lys Glu
 210 215 220

Tyr Leu Lys Thr Phe Asn Ala Met Glu Leu Glu Val Asn Pro Lys Ser
 225 230 235 240

Val Gly Leu Pro Gly Ser Pro Thr Ile Val Tyr Lys Val Glu Lys Ile
 245 250 255

Pro Arg Ala Lys Ala Asn Arg Lys Ala Asp Val Val Asp Gly Ser Asn
 260 265 270

Gln Asp Ser Leu Arg Gln Val Ala Arg Arg Ile His Asp Val Leu Gly
 275 280 285

Gly Val Val Ile Lys
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<210> 26

<211> 1284

<212> DNA

<213> Crenarchaeote

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<210> 27

<211> 427

<212> PRT

<213> Crenarchaeote

<400> 27

Met Thr Ser Ser Leu Ser Ala Ile Pro Asp Ala Lys Leu Asp Glu Arg
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Pro Asn Gln Asn Ala His Val Asn Asp Asn Pro Glu Lys Glu Arg Gly
20 25 30

Asp Asn Asn Arg His Leu Tyr Val Val Ile Glu Gln Glu Glu Gly Thr
35 40 45

Ile Leu Pro Val Ser Phe Glu Met Leu Gly Glu Ala Arg Arg Leu Met
50 55 60

Asp Asp Phe Asn His Lys Tyr Lys Pro Glu Glu Lys Val Val Ala Ile
65 70 75 80

Ile Leu Gly His Asn Ile Lys His Leu Cys Gln Glu Leu Ile His His
85 90 95

Gly Ala Asp Ala Val Ile Tyr Ala Asp His Pro Glu Leu Arg His Pro
100 105 110

Arg Asn Leu Leu Tyr Thr Lys Val Val Cys Gln Ile Ala Thr Asp Lys
115 120 125

Glu Ser Ala Ala Arg Ile Trp Pro Ser Asn Pro Asp Phe Asn Arg Pro
 130 135 140

Arg Tyr Met Phe Phe Ser Ala Asp Asp Thr Gly Arg His Leu Ser Ser
 145 150 155 160

Thr Val Leu Ala Glu Leu Gln Ser Gly Leu Ala Ser Asp Ile Asn Lys
 165 170 175

Leu Val Ile Asn Asp Leu Glu Ile Arg His Glu His Lys Thr Lys Gly
 180 185 190

Lys Pro Ile Val Tyr Glu Lys Thr Leu Glu Met Tyr Arg Pro Asp Phe
 195 200 205

Ser Gly Phe Leu Trp Thr Thr Ile Leu Cys Leu Asp Asn Ile Asn Pro
 210 215 220

Glu Asn Arg Arg Lys Phe His Pro Gln Ala Cys Ser Ile Ile Pro Gly
 225 230 235 240

Val Phe Pro Gln Met Glu Gly Asp Thr Asp Arg Lys Gly Thr Ile Ile
 245 250 255

Glu Phe Ser Pro Thr Ile Ala Gln Glu Asp Leu Arg Ile Lys Ile Ile
 260 265 270

Asn Arg Arg Val Ile Lys Ser Lys Val Asp Phe Ser Asn Lys Lys Ile
 275 280 285

Ile Val Ser Phe Gly Arg Gly Ile Lys Glu Ser Pro Glu Gln Asn Ile
 290 295 300

Lys Leu Ile Glu Asn Leu Ala Lys Glu Ile Glu Ala Glu Ile Gly Ile
 305 310 315 320

Ser Leu Pro Ile Ser Lys Lys Pro Tyr Pro Ile Ser Glu Ser Leu Ser
 325 330 335

Ser Thr Tyr Met Ile Pro Asp Arg Val Ile Gly Thr Ser Gly Arg Lys
 340 345 350

Val Asn Pro Gln Val Tyr Phe Ala Ile Gly Ile Ser Gly Ala Val Gln
 355 360 365

His Ile Ala Gly Met Lys Glu Ser Glu Phe Val Ile Ser Ile Asn Pro
 370 375 380

Asp Ser Glu Ala Pro Ile Ile Asp Glu Ser Asp Val Leu Ile Lys Gly
 385 390 395 400

Lys Ile Glu Gln Val Leu Pro Leu Leu Ile Asn Glu Leu Lys Lys Tyr
 405 410 415

Lys Glu Arg Leu Gln Ile Pro Gln Glu Ile Glu
 420 425

<210> 28

<211> 1878

<212> DNA

<213> Crenarchaeote

<400> 28

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<210> 29

<211> 625

<212> PRT

<213> Crenarchaeote

<400> 29

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 20 25 30

Leu Ile Glu Ala Gly Lys Lys Ile Gly Thr Lys Asn Val Ser Gly Gly
 35 40 45

Ile Leu Tyr Ser Lys Lys Thr Ala Thr Gly Lys Val His Asn Val Glu
 50 55 60

Asp Val Phe Asp Asn Phe Leu Ala Asp Ala Pro Leu Glu Arg Lys Ile
 65 70 75 80

Ile Lys Tyr Met Leu His Ala Val Ser Arg Glu Lys Ala Phe Ser Leu
 85 90 95

Asp Leu Thr Leu Ala His Asp Tyr Gln Thr Asn Phe Gly Tyr Thr Val
 100 105 110

Leu Leu Asn Lys Leu Leu Ser Trp Phe Ala Arg Glu Ala Ser Gln Ser
 115 120 125

Ala Glu Lys Leu Gly Gly Gly Ile Ile Thr Gly Val His Leu Arg Ser
 130 135 140

Ile Ile Trp Lys Asp Asp Ser Thr Ile Ile Ile Glu Thr Asp Glu Leu
 145 150 155 160

Glu Pro Phe Gln Val Lys Ala Val Ile Ala Ala Asp Gly Val Asn Ser
 165 170 175

Glu Val Ala Gln Ile Thr Gly Ala Arg Ser Lys Phe Thr Pro Ser Asp
 180 185 190

Leu Tyr Gln Gly Val Lys Val Val Ala Lys Leu Pro Glu Gly Leu Leu
 195 200 205

Glu Glu Arg Phe Gly Val Ser Glu Asn Glu Gly Ala Ala His Leu Phe
 210 215 220

Ser Gly Asp Ile Thr Leu Asn His Ile Gly Gly Gly Phe Leu Tyr Thr
 225 230 235 240

Asn Arg Asp Thr Ile Ser Ile Gly Ala Val Tyr His Tyr Asp Ser Leu
245 250 255

Ile Glu Lys Pro Thr Glu Pro Asn Ala Leu Val Asn Ala Leu Leu Ser
260 265 270

Asn Pro Phe Val Met Glu Leu Ile Lys Asp Glu Val Pro Arg Ile Lys
275 280 285

Glu Asp Tyr Arg Asp Leu Ser Lys Asp Glu Glu Leu Arg Ile Arg Phe
290 295 300

Lys Ser Asn Lys Leu Ile Lys Ser Trp Asn Asp Leu His His Thr Tyr
305 310 315 320

Tyr Ser Pro Ser Ala Val Ala Glu Leu Val Ala Gln Gly Lys Tyr Lys
325 330 335

Ser Arg Glu Glu Ile Lys Asp Lys Ile Asp Ser Leu Tyr Asn Glu Leu
340 345 350

Val Thr Lys Tyr Asn Thr Glu Phe Glu Thr Asn Tyr Val Glu Leu Glu
355 360 365

Tyr Ser Ala Lys Leu Val Pro Asp Gly Lys Arg Cys Arg Met Lys Lys
370 375 380

Pro Tyr Phe Lys Asn Ile Leu Phe Val Gly Asp Ala Ala Gly Arg Gly
385 390 395 400

Ile Phe Leu Gly Pro Arg Ile Glu Gly Leu Asn Val Gly Ile Asp Asp
405 410 415

Ala Val Arg Ala Ala Glu Ala Val Ser Lys Ser Ile Asp Gln Asn Asn
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Phe Gln Phe Asp Asn Ile Gly Glu Arg Tyr Thr Lys Ser Val Asp Glu
435 440 445

Ser Pro Tyr Thr Ala Asp Met Ser Arg Ile Asp Ala Asn Tyr Leu Lys
450 455 460

Ala Val Leu Asp Cys Thr Lys Lys Val Pro Lys Asn Thr Leu Gly Phe
465 470 475 480

Lys Tyr Gly Ser Ile Val Lys Leu Met Ser Asn Ser Thr Phe Arg Asn
485 490 495

Val Ser Ile Gly Ile Ala Asn Ser Ile Gly Tyr Lys Arg Leu Leu Pro
500 505 510

Val Ile Glu Ser Asp Lys Thr Tyr Asn Gln Ile Pro Ile Glu Ile Ala
515 520 525

Glu Arg Asn Gly Lys Asp Leu Arg Lys Ser Tyr Ser Ile Glu Ile Pro
530 535 540

Thr Ile Ala Glu Arg Ile Ala Asn Leu Asn Tyr Asn Asp Asp Ser Leu
545 550 555 560

Ser His Ile Lys Val Leu Asn Ser Gln Ser Asp Phe Met Lys Lys Met
565 570 575

Val Gln Leu Cys Pro Thr Lys Cys Tyr Ser Ile Glu Asn Glu Arg Ile
580 585 590

Met Leu Gln His Glu Gly Cys Ile Glu Cys Gly Thr Cys Ala Arg Glu
595 600 605

Thr Glu Trp Arg His Pro Arg Gly Glu Lys Gly Ile Ile Tyr Asn Tyr
610 615 620

Gly
625

<210> 30

<211> 2238

<212> DNA

<213> Crenarchaeote

<400> 30

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gtcctctatg accagagtgt cctgttccag cagcttgacc aaaacgacaa agtcaaaagc     1140
gagttcataa acgtggctgc gcatgagttg cgaacgccaa tcatgcccac cctaaacggt     1200
gtggagatac tggaggagaa gcttggcgaa agaaaaacag agtttcagcg ggagcttgac     1260
atgataacaa gaaacgcgtc ccggtgcag aaccttgccg aaagcattct gcaggtgagc     1320
agaatcgaag gcggaagctt tagcctggac atccaaaaaa atgtggatat ccacaacctg     1380
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atttcccagg tgatagagga cattgagaaa aaatacgcct acaaggagaa ggcaaacaag 1440
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 ggcggcgcaa aggcagaagg ggtaaaggca gcggcaggag caaaacaggc gcaaaaagag 1560
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 ggcagagtgg atagcagcag cgacagcgac aacgggtggtg gtgacaatgg tggtgaccac 1860
 atcgggaggc agaaagaagg cgcggtgcta gtcacagtgc aggacaccgg ggttgggctc 1920
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<210> 31

<211> 745

<212> PRT

<213> Crenarchaeote

<400> 31

Met Glu Gly Ser Ser Leu Ile His Asn Asn Asn Leu His Asp Ile Gly
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Thr Asn Asn Asp Asn Val Cys Glu Asn Lys Tyr Ile Asp Ala Asn Pro
 20 25 30

Gly Leu Ser His Lys Arg Gly Phe Gly Asp Val Asn Asn Asn Glu Asn
 35 40 45

Asp Asn Asp Gly Gly Val Asp Ala Gly Ala Pro Thr Thr Lys Val Tyr
 50 55 60

Tyr Gly Pro Glu Asn Ala Asn Asn Ala Ile Leu Arg Phe Ile Asp Arg
 65 70 75 80

Ala Asn Val Lys Ile Asp Ser Cys Ile Asn Ser Val Ala Pro Ser Val
 85 90 95

Met Ile Gly Val Asp Ala Ile Arg Glu Lys Arg Val Asp Ala Val Lys
 100 105 110

Asn Arg Gly Leu Lys Leu Arg Tyr Val Thr Glu Ile Thr Lys Asp Asn
 115 120 125

Val Gly Tyr Val Lys Glu Met Leu Ser Phe Ser Glu Ile Arg His Leu
 130 135 140

Asp Gly Leu Lys Gly Asn Phe Glu Val Ala Asp Gln Arg Glu Tyr Val
 145 150 155 160

Ala Val Ala Thr Leu His Ala Ala Gln Ser Ile Pro Gln Leu Leu Phe
 165 170 175

Ser Asn Leu Pro Glu Ile Ala Glu Gln Gln Gln Phe Val Phe Asp Ser
 180 185 190

Phe Trp Gly Arg Ala Leu Pro Ala Glu His Arg Ile Lys Glu Leu Glu
 195 200 205

Asp Gly Val Val Met Pro Val Ser Leu Val Phe Ser Asn Tyr Lys Asp
 210 215 220

Ala Val Gln Arg Glu Phe Glu Met Ile Ser Arg Ala Lys Arg Glu Ile
 225 230 235 240

Leu Ile Met Tyr Ser Thr Val Asn Ala Phe His Leu Gln Glu Lys Gly
 245 250 255

Gly Thr Leu Gln Leu Leu Lys Glu Met Val Glu Gln Asn Asp Ser Leu
 260 265 270

Arg Ile Asn Ile Leu Thr Pro Met Asp Ala Ser Val Arg Glu Ser Leu
275 280 285

Ser Leu Arg Leu Leu Thr Lys Tyr Arg Pro Asn Ile Gln Val Gln Asp
290 295 300

Ile Ala Pro Ser Ile Gly Ile Lys Ile Lys Thr Leu Val Val Asp Arg
305 310 315 320

Lys Glu Ser Leu Val Met Glu Leu Ile His Ala Arg Glu Glu Val Ala
325 330 335

Thr Ala Ala Ile Gly Phe Ser Ile Tyr Ser Asn Ser Glu Pro Thr Val
340 345 350

Leu Ser Tyr Ser Ser Ile Phe Glu Val Leu Tyr Asp Gln Ser Val Leu
355 360 365

Phe Gln Gln Leu Asp Gln Asn Asp Lys Val Lys Ser Glu Phe Ile Asn
370 375 380

Val Ala Ala His Glu Leu Arg Thr Pro Ile Met Pro Ile Leu Asn Gly
385 390 395 400

Val Glu Ile Leu Glu Glu Lys Leu Gly Glu Arg Lys Thr Glu Phe Gln
405 410 415

Arg Glu Leu Asp Met Ile Thr Arg Asn Ala Ser Arg Leu Gln Asn Leu
420 425 430

Ala Glu Ser Ile Leu Gln Val Ser Arg Ile Glu Ser Gly Ser Phe Ser
435 440 445

Leu Asp Ile Gln Lys Asn Val Asp Ile His Asn Leu Ile Ser Gln Val
450 455 460

Ile Glu Asp Ile Glu Lys Lys Tyr Ala Tyr Lys Glu Lys Ala Asn Lys
465 470 475 480

Val Ala Ile Val Phe Leu Pro Ser Asp Gly Asn Arg Asn Gly Gly Tyr
485 490 495

Ser Arg Gly Gly Gly Gly Ala Lys Ala Glu Gly Val Lys Ala Ala Ala
500 505 510

Gly Ala Lys Gln Ala Gln Lys Glu Thr Gln Gln Lys Glu Gln Trp Val
515 520 525

Glu Pro Val Asn Gly Pro Asn His Leu Leu Tyr Val Asp Cys Asp Pro
530 535 540

Gln Lys Ile Ser Gln Val Val Phe Asn Leu Leu Asp Asn Ala Met Lys
545 550 555 560

Phe Thr Asn Asp Gly Lys Ile Val Val Ser Thr Ala Val Met Gly Glu
565 570 575

Ser Ser Pro Phe Thr Ser Thr Ser Gln Glu Ser Asp Thr Ser Asn Thr
580 585 590

Ala Thr Ala Gly Lys Gly Asn Gly Gly Arg Val Asp Ser Ser Ser Asp
595 600 605

Ser Asp Asn Gly Gly Gly Asp Asn Gly Gly Asp His Ile Gly Arg Gln
610 615 620

Lys Glu Gly Ala Val Leu Val Thr Val Gln Asp Thr Gly Val Gly Leu
625 630 635 640

Asn Ser Lys Ile Arg Asp Gln Leu Phe Gln Lys Phe Val Thr Lys Ser
645 650 655

Asn Gln Gly Thr Gly Leu Gly Leu Tyr Leu Ser Arg Lys Ile Val Glu
660 665 670

Glu His Gly Gly Lys Ile Trp Phe Glu Glu Thr Asn Ser Lys Gly Gly
675 680 685

Asn Ser Ser Ser Arg Asn Asn Thr Lys Asp Lys Asp Glu Gly Ile Asp
690 695 700

Glu Ile Leu His His Leu Gly Ser Glu Gly Lys Ile Gly Ala Thr Phe
705 710 715 720

Lys Phe Val Ile Pro Val Ser Leu Pro Ser His Met Pro Thr Lys Asp
725 730 735

Met Pro Glu Lys Asn Asp Glu Gly Lys
740 745

<210> 32

<211> 519

<212> DNA

<213> Crenarchaeote

<400> 32

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acaagtcact tgattattaa aatatatcca catatggata acaatacaag gatgagttct	180
ttagcaatcg agtttttttt atcccttttt tcaataacgt tacttttctaa aagaatatac	240
caaccagtga aatcaaagtc atatacctac catgacaagc atccatttca gtacaagatg	300
gaggattatg caaaccacaa caaaattgta gactataaaa actgcttact tttttttcaa	360
gtatcgatgt tacaaaaaaaa taaaataatt aggattcggg ttccagggtt gttttataca	420
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<210> 33

<211> 172

<212> PRT

<213> Crenarchaeote

<400> 33

Met Gln Ser Ser His Leu Ser Lys Ile Ile Thr Ile Cys Arg Met Pro
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Ser Leu His Leu Val Ala Tyr Gly Leu Ile Leu Asp Ile Phe Glu Ser
20 25 30

Pro Ile Thr Arg Leu Asn Gly Arg Thr Ser His Leu Ile Ile Lys Ile
 35 40 45

Tyr Pro His Met Asp Asn Asn Thr Arg Met Ser Ser Leu Ala Ile Glu
 50 55 60

Phe Phe Leu Ser Leu Phe Ser Ile Thr Leu Leu Ser Lys Arg Ile Tyr
 65 70 75 80

Gln Pro Val Lys Ser Lys Ser Tyr Thr Tyr His Asp Lys His Pro Phe
 85 90 95

Gln Tyr Lys Met Glu Asp Tyr Ala Asn His Asn Lys Ile Val Asp Tyr
 100 105 110

Lys Asn Cys Leu Leu Phe Phe Gln Val Ser Met Leu Gln Lys Asn Lys
 115 120 125

Ile Ile Arg Ile Arg Val Pro Gly Leu Phe Tyr Thr Gly Gly Trp Ile
 130 135 140

Ser Leu Thr Leu Lys Phe Leu Ile Ser Thr Ser Phe Ala Pro Ser His
 145 150 155 160

Leu Lys Val Ala Met Gly Pro Pro Gln Asp Ile Ile
 165 170

<210> 34

<211> 1008

<212> DNA

<213> Crenarchaeote

<400> 34

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aacacagttg cgaccctaaa cccctctgca gatttaacac ctggtgccac atacactgcc	120
agaataacaa ccggtgctac ggatgcgact ggtgtcccat tggctgccga caaggtatgg	180
acgttctctg ttgccccctt tggaggcgga gggacattag accagtttgg gataaccag	240
atttaccocg ataaggcggg tgggtggagaa aaatggttta tgaatatgca aaaccggaac	300

aatgatccac gaacaaaccc acctgacatg gacctaaacc cagatggcag ttggaacgtt 360
aatgacgac aggtcagata taacgtgttt acatcatcag ggtaccatcc agaggatatt 420
gagacttacg atcactcggg actcgcaaca caaggataca tgcagtatcc aatgattgg 480
aagaatgtgg aaatgacggg tattgtaagg gttaatagtg gagatgattc. tgaaaatttc 540
gcttggtatg acaggggtgg taggcactat gatggcgaag gatgagaggg ctcagcatat 600
aaagcagatc tattctatga tggaaggggtt aggcttgcaa aagagcagtg gcatgtctcc 660
tatgtgtttt caagcactac cacaccctca ccttcggcgt ctagttttga tagattcatt 720
ggatttaaag ccatgatcta taaccaccaa ttggctggtg gtgagaccgt agtcaccact 780
gaaatatggg tagacagaaa cccggaatgc ccgactctga agaacaattg gcaaaaggta 840
tacacattca ctgactcagg tgggttttga aatgatggtg aagagtgtgg tggtagagccg 900
gatcagatta tatcctgggg aggccccatt gctactttca ggtgggatgg tgcaaatgat 960
gtggatatca aaaactttag tgtgagggaa atccagccac ctgtataa 1008

<210> 35

<211> 335

<212> PRT

<213> Crenarchaeote

<400> 35

Thr Leu Leu Thr Ser Gly Gly Ala Pro Val Pro Ala Thr Val Ser Met
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Asn Ser Pro Thr Asn Thr Val Ala Thr Leu Asn Pro Ser Ala Asp Leu
20 25 30

Thr Pro Gly Ala Thr Tyr Thr Ala Arg Ile Thr Thr Gly Ala Thr Asp
35 40 45

Ala Thr Gly Val Pro Leu Ala Ala Asp Lys Val Trp Thr Phe Ser Val
50 55 60

Ala Pro Ser Gly Gly Gly Gly Thr Leu Asp Gln Phe Gly Ile Thr Gln
65 70 75 80

Ile Tyr Pro Asp Lys Ala Gly Gly Gly Glu Lys Trp Phe Met Asn Met
85 90 95

Gln Asn Pro Asn Asn Asp Pro Arg Thr Asn Pro Pro Asp Met Asp Leu
100 105 110

Asn Pro Asp Gly Ser Trp Asn Val Asn Asp Asp Gln Val Arg Tyr Asn
115 120 125

Val Phe Thr Ser Ser Gly Tyr His Pro Glu Asp Ile Glu Thr Tyr Asp
130 135 140

His Ser Val Leu Ala Thr Gln Gly Tyr Met Gln Tyr Pro Asn Asp Trp
145 150 155 160

Lys Asn Val Glu Met Thr Gly Ile Val Arg Val Asn Ser Gly Asp Asp
165 170 175

Ser Glu Asn Phe Ala Trp Tyr Asp Arg Gly Gly Arg His Tyr Asp Gly
180 185 190

Glu Gly Cys Glu Gly Ser Ala Tyr Lys Ala Asp Leu Phe Tyr Asp Gly
195 200 205

Arg Val Arg Leu Ala Lys Glu Gln Trp His Val Ser Tyr Val Phe Ser
210 215 220

Ser Thr Thr Thr Pro Ser Pro Ser Ala Ser Ser Phe Asp Arg Phe Ile
225 230 235 240

Gly Phe Lys Ala Met Ile Tyr Asn His Gln Leu Ala Gly Gly Glu Thr
245 250 255

Val Val Thr Thr Glu Ile Trp Val Asp Arg Asn Pro Asp Ser Pro Thr
260 265 270

Leu Lys Asn Asn Trp Gln Lys Val Tyr Thr Phe Thr Asp Ser Gly Gly
275 280 285

Phe Gly Asn Asp Gly Glu Glu Cys Gly Gly Glu Pro Asp Gln Ile Ile
290 295 300

Ser Trp Gly Gly Pro Ile Ala Thr Phe Arg Trp Asp Gly Ala Asn Asp
305 310 315 320

Val Asp Ile Lys Asn Phe Ser Val Arg Glu Ile Gln Pro Pro Val
325 330 335